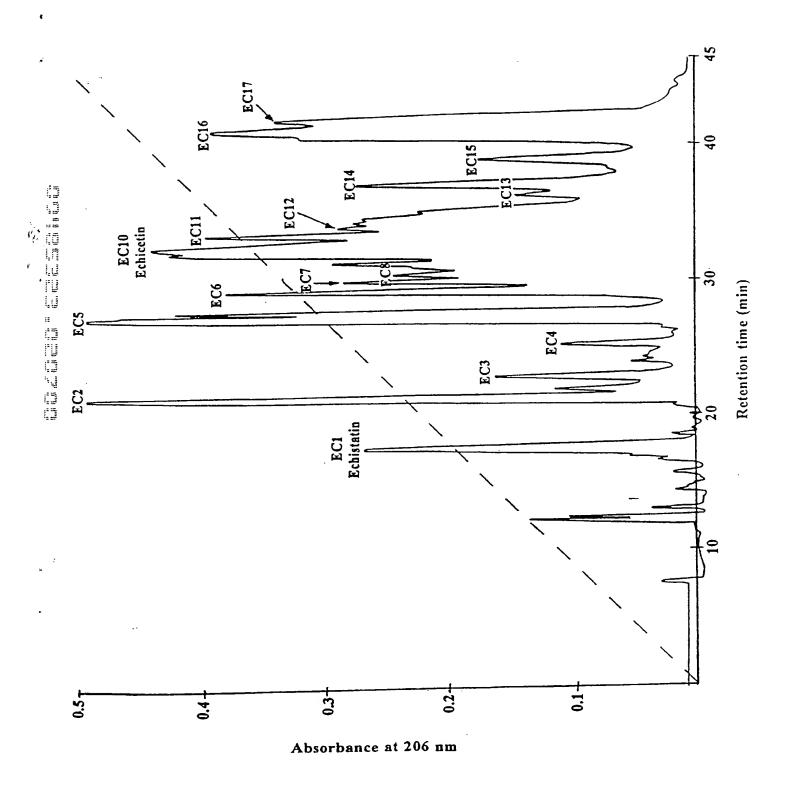
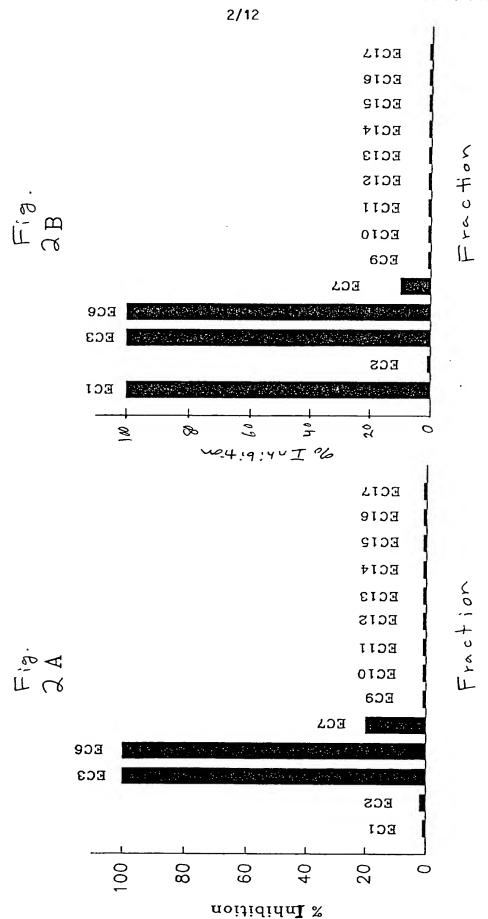
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Fig. 1







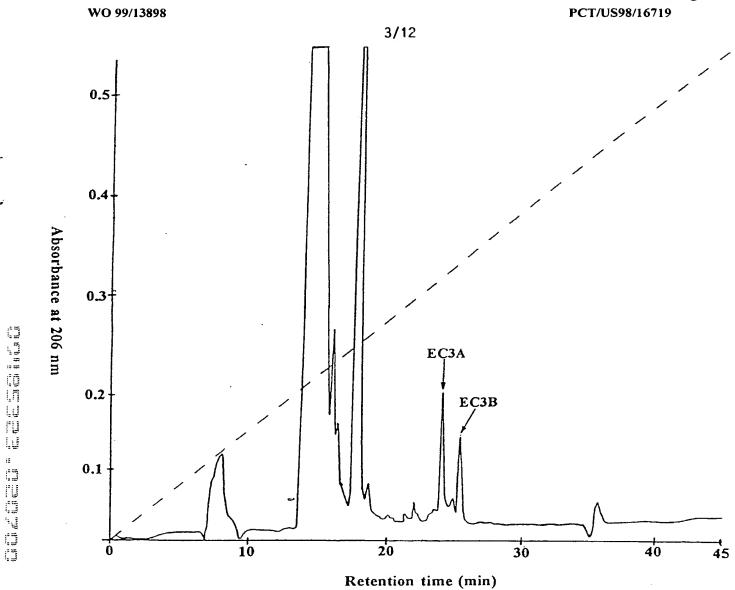


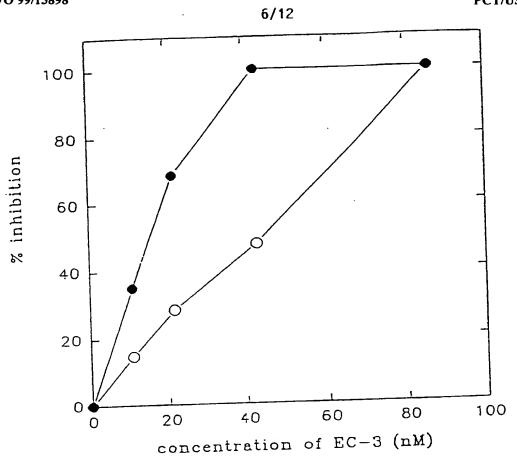
Fig. 3

	Fig. 4A		
60 TPDCPRNRYKGKED	QQ	K4	
50 DDYCSG1		1	
40 STVCKRA <u>VGD</u> DV	 	Kla   K Klb V -    Kla  N K2 I -	
30 PCCRNCYFLRA	terminal sequencing	K Kla N K	
20 GEHCISG	- N-termi	<u> </u>	
1 50 60 60 NSVHPCCDPVKCEPREGEHCISGPCCRNCYFLRAGTVCKRAVGDDVDDYCSGITPDCPRNRYKGKED		SA	

	F:3.4B	
10 20 30 40 50 60 NSVHPCCDPVKCEPREGEHCISGPCCRNCKFLNAGTICKRAMLDGLNDYCTGISTDCPRNRYKGKED T	1 sequencing	K10b S
50 AMLDGLNDYCTC		( K10b
30 CKFLNAGTICKR	sequencing	<del></del>
20 BEHCISGPCCRN	N-terminal sequencin CNBr-1	K4
10 CCDPVKCEPREG T	N-terminal sequencing	T K4
1 NSVHP(		

	L	ſ
	<i></i>	

	CI'GKSCUDPRN PWNG	CNGKTCDCPRNPHKGPAT	FPDDRCTGLSNDCPRWNDL	CTGQSADCPRYH	CSGITPDOPRNRYKGKED	CTGKSSDCPRNRYKGKED	drgissdernpykd
	MNDO	DMDD	FPDD]	MPDD	DVDD	GIND	ОДОМО
, ,	<u> </u>	RGD	RGD	RGD	VGD	MLD	<u>VGD</u>
	¥ 	OKRA	GRIA	GRIP	d K R A	CKRA	dkra
The state of the s	VEEF CAIGE CHECKEN THEORY	EGESGPCGRNCKFLKEGTICKRA RGD DMDDYCNGKTCDCPRNPHKGPAT	GEECDCSSPSNPCCDAATCKLRPGAQCADGLCCDQCRFKKKTGICRIA RGD	GKECDCSSPENPCCDDATCKLRPGACCEGLCCEQCKFSRAGKICRIP RGD MPDDRCTGQSADCPRYH	NSVHPCCDPVKGEPREGEHCISGPCCRNCKFLRAGTVCKRA <u>VGD</u> DVDDYCSGITPDCPRNRYKGKED	NSVHPCCDPVKCEPREGEHCISGPCCRNCKFLNAGTICKRA <u>MLD</u> GLNDYCTGKSSDCPRNRYKGKED	nsgnpccppvidpprrgehdvsgkcdrndkflragivdkra <u>vgd</u> dmddydigissddprnpykd
Eristostatin	Echistatin	Flavoridin	Kistrin	EC3A	EC3B	Le3 (disintegrin domain)	



F.g. 6



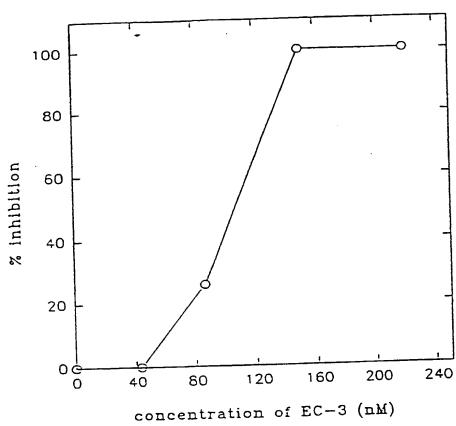
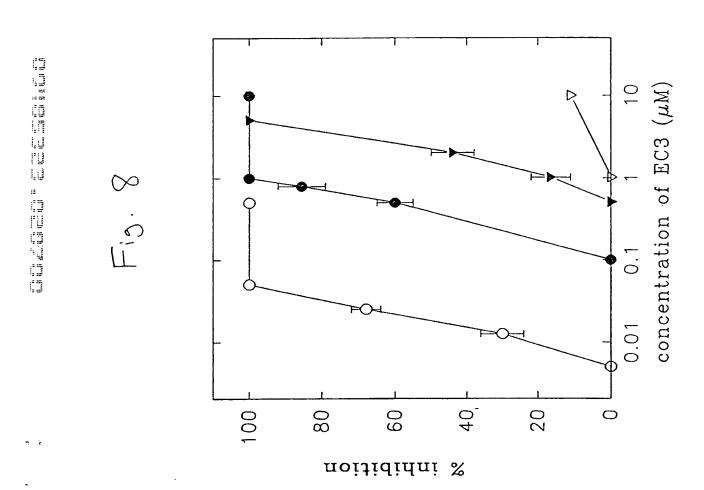
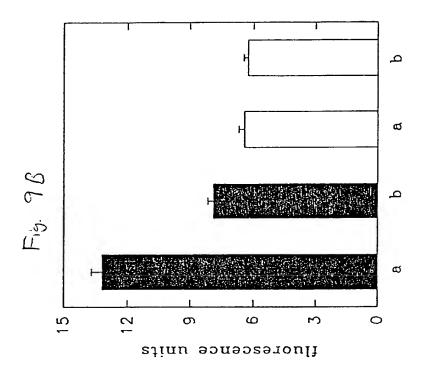
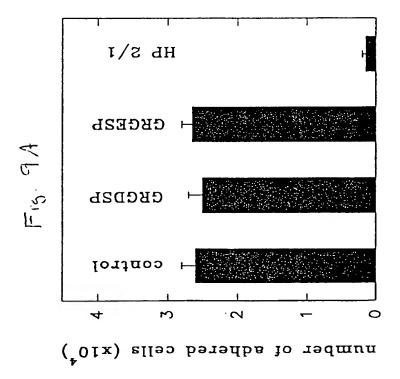


Fig. 7





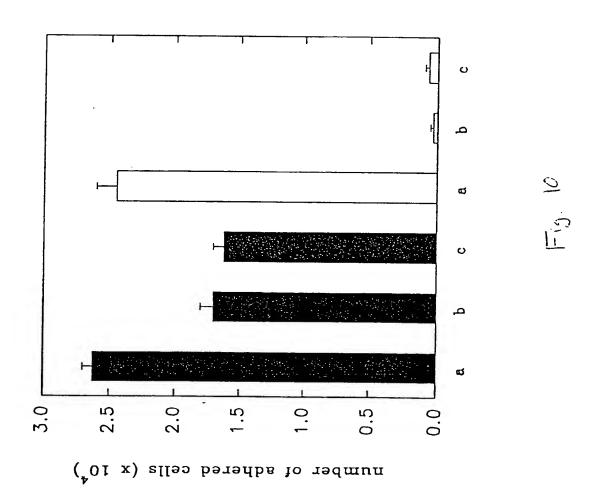


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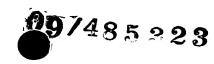
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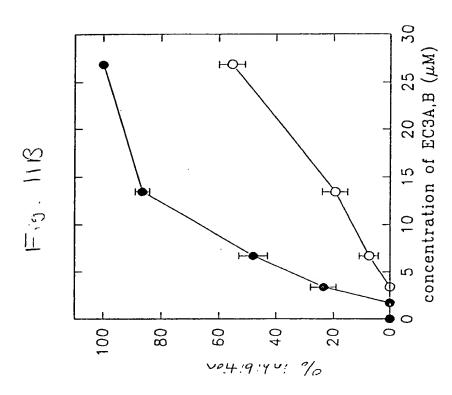


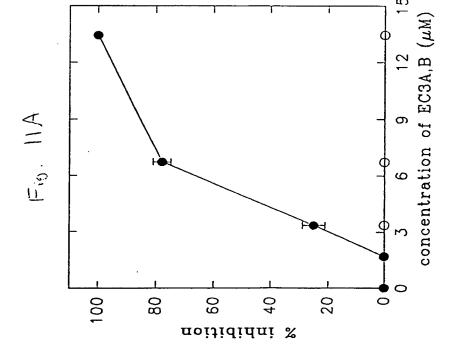
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